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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/533,466

DATE: 03/29/2001
TIME: 16:16:47

ENTERED

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Output Set: N:\CRF3\03292001\I533466.raw

3 <110> APPLICANT: COLLART, FRANK R.
4 HUBERMAN, ELIEZER
5 JOACIMIAK, ANDRZEJ
6 ZHANG, RONGGUANG
7 WESTBROOK, EDWIN M.
9 <120> TITLE OF INVENTION: USE OF CRYSTAL STRUCTURE OF BACTERIAL IMP DEHYDROGENASE
10 TO DESIGN INHIBITORS OF BACTERIAL GROWTH
12 <130> FILE REFERENCE: 21416/90042
14 <140> CURRENT APPLICATION NUMBER: 09/533,466
15 <141> CURRENT FILING DATE: 2000-03-23
17 <160> NUMBER OF SEQ ID NOS: 23
19 <170> SOFTWARE: PatentIn Ver. 2.1
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44 <213> ORGANISM: Escherichia coli
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53 <212> TYPE: PRT
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63 <212> TYPE: PRT
64 <213> ORGANISM: Mycobacterium tuberculosis
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68 1 5 10 15

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140 Asp Lys Leu Val Pro Glu Gly
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147 <213> ORGANISM: Bacillus subtilis
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160 <213> ORGANISM: Escherichia coli
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167      20
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171 <211> LENGTH: 21
172 <212> TYPE: PRT
173 <213> ORGANISM: Bacillus subtilis
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176 Met Glu Lys Gly Ser Lys Asp Arg Tyr Phe Gln Glu Glu Asn Lys Lys
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184 <211> LENGTH: 30
185 <212> TYPE: PRT
186 <213> ORGANISM: Mycobacterium tuberculosis
188 <400> SEQUENCE: 16
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193      20      25      30
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198 <212> TYPE: PRT
199 <213> ORGANISM: Homo sapiens
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205 Lys Ile Lys Val Ala Gln Gly
206      20
209 <210> SEQ ID NO: 18
210 <211> LENGTH: 23

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224 <212> TYPE: PRT
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232           20           25
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245           20           25
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249 <211> LENGTH: 21
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251 <213> ORGANISM: Drosophila melanogaster
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275 <211> LENGTH: 477
276 <212> TYPE: PRT
277 <213> ORGANISM: Artificial Sequence
279 <220> FEATURE:
280 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic

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281      polypeptide
283 <220> FEATURE:
284 <221> NAME/KEY: MOD_RES
285 <222> LOCATION: (1)..(477)
286 <223> OTHER INFORMATION: "Xaa" represents selenomethionine
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292 Val Leu Leu Ile Pro Ala Glu Ser His Val Leu Pro Asn Glu Val Asp
293      20      25      30
295 Leu Lys Thr Lys Leu Ala Asp Asn Leu Thr Leu Asn Ile Pro Ile Ile
296      35      40      45
W--> 298 Thr Ala Ala Xaa Asp Thr Val Thr Gly Ser Lys Xaa Ala Ile Ala Ile
299      50      55      60
W--> 301 Ala Arg Ala Gly Gly Leu Gly Val Ile His Lys Asn Xaa Ser Ile Thr
302  65      70      75      80
304 Glu Gln Ala Glu Glu Val Arg Lys Val Lys Arg Ser Glu Asn Gly Val
305      85      90      95
307 Ile Ile Asp Pro Phe Phe Leu Thr Pro Glu His Lys Val Ser Glu Ala
308      100     105     110
W--> 310 Glu Glu Leu Xaa Gln Arg Tyr Arg Ile Ser Gly Val Pro Ile Val Glu
311      115     120     125
W--> 313 Thr Leu Ala Asn Arg Lys Leu Val Gly Ile Ile Thr Asn Arg Asp Xaa
314      130     135     140
W--> 316 Arg Phe Ile Ser Asp Tyr Asn Ala Pro Ile Ser Glu His Xaa Thr Ser
317 145     150     155     160
319 Glu His Leu Val Thr Ala Ala Val Gly Thr Asp Leu Glu Thr Ala Glu
320      165     170     175
322 Arg Ile Leu His Glu His Arg Ile Glu Lys Leu Pro Leu Val Asp Asn
323      180     185     190
325 Ser Gly Arg Leu Ser Gly Leu Ile Thr Ile Lys Asp Ile Glu Lys Val
326      195     200     205
328 Ile Glu Phe Pro His Ala Ala Lys Asp Glu Phe Gly Arg Leu Leu Val
329      210     215     220
331 Ala Ala Ala Val Gly Val Thr Ser Asp Thr Phe Glu Arg Ala Glu Ala
332 225     230     235     240
334 Leu Phe Glu Ala Gly Ala Asp Ala Ile Val Ile Asp Thr Ala His Gly
335      245     250     255
337 His Ser Ala Gly Val Leu Arg Lys Ile Ala Glu Ile Arg Ala His Phe
338      260     265     270
340 Pro Asn Arg Thr Leu Ile Ala Gly Asn Ile Ala Thr Ala Glu Gly Ala
341      275     280     285
343 Arg Ala Leu Tyr Asp Ala Gly Val Asp Val Val Lys Val Gly Ile Gly
344      290     295     300
346 Pro Gly Ser Ile Cys Thr Thr Arg Val Val Ala Gly Val Gly Val Pro
347 305     310     315     320
349 Gln Val Thr Ala Ile Tyr Asp Ala Ala Ala Val Ala Arg Glu Tyr Gly
350      325     330     335
352 Lys Thr Ile Ile Ala Asp Gly Gly Ile Lys Tyr Ser Gly Asp Ile Val

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VERIFICATION SUMMARY
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DATE: 03/29/2001
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Input Set : A:\21416942.app
Output Set: N:\CRF3\03292001\I533466.raw

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